

Figure 1

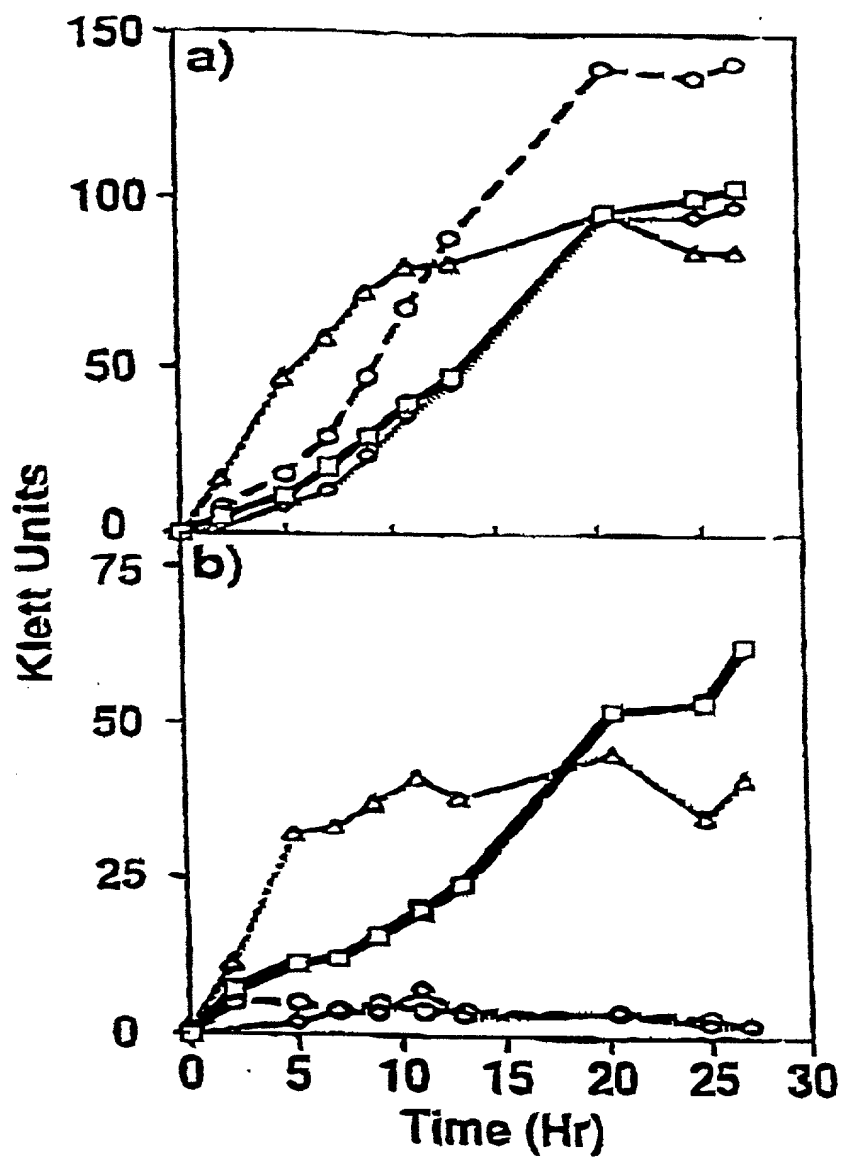


Figure 2

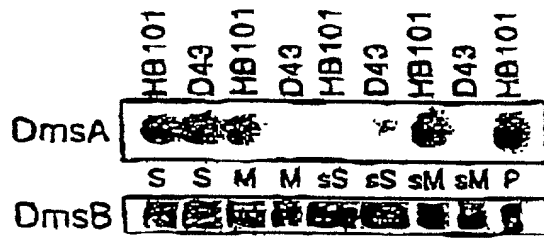


Figure 3

a)

b)

c)

1 2 3 4 5 6 1 2 1 2

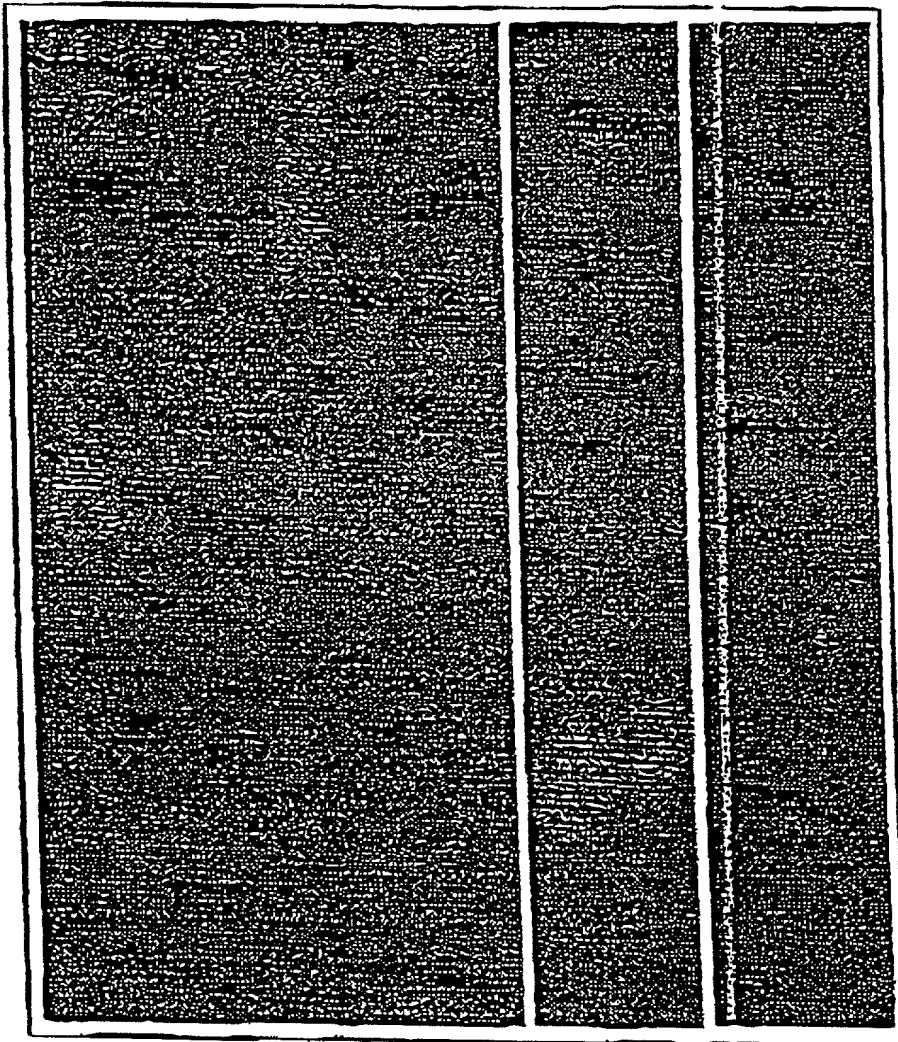
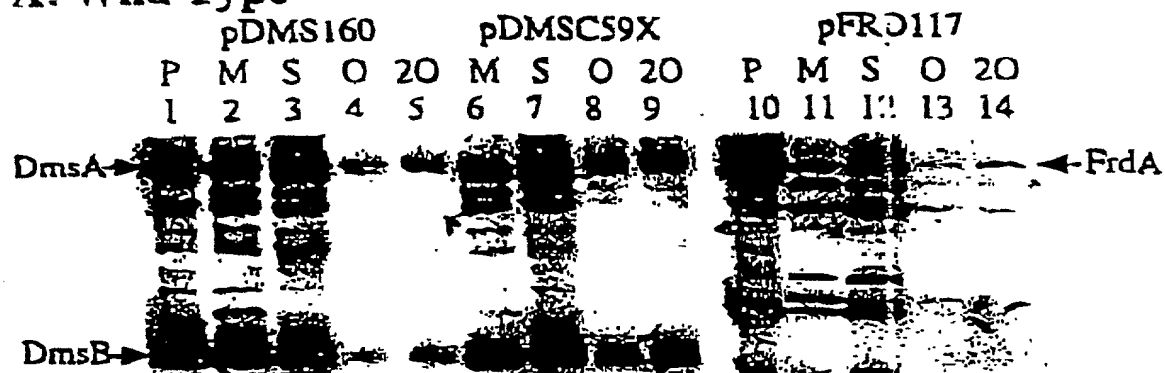


Figure 4

A: Wild Type



B: D43

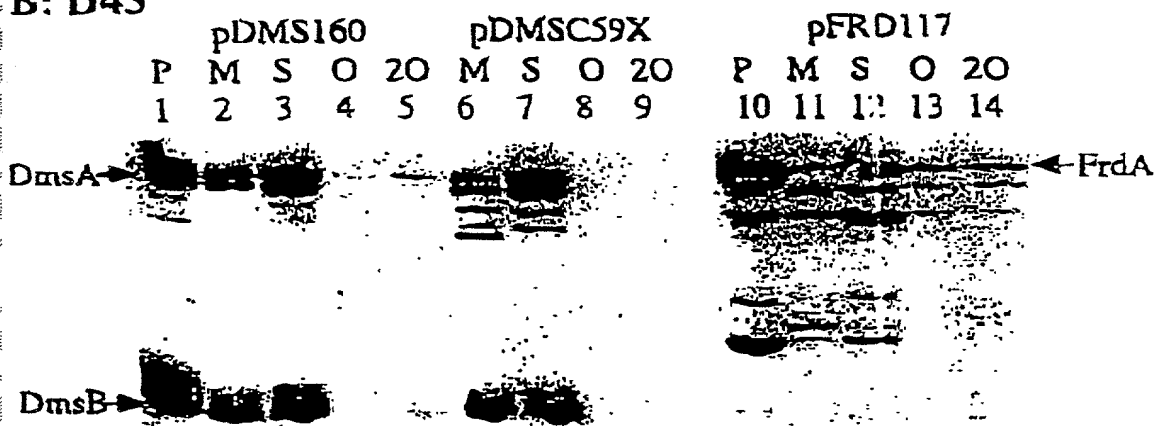


Figure 5

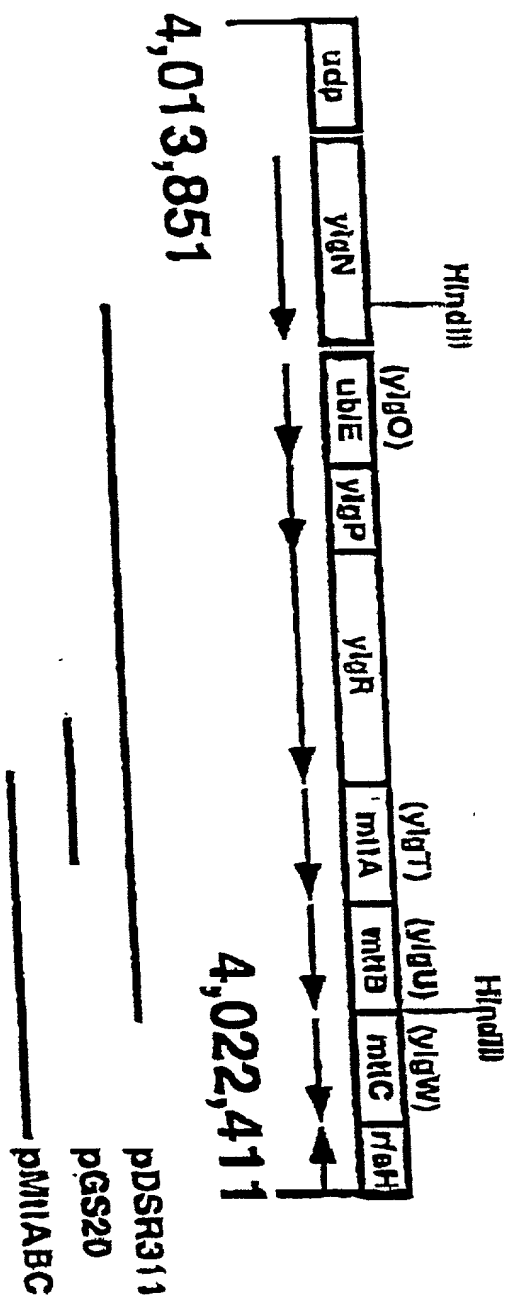
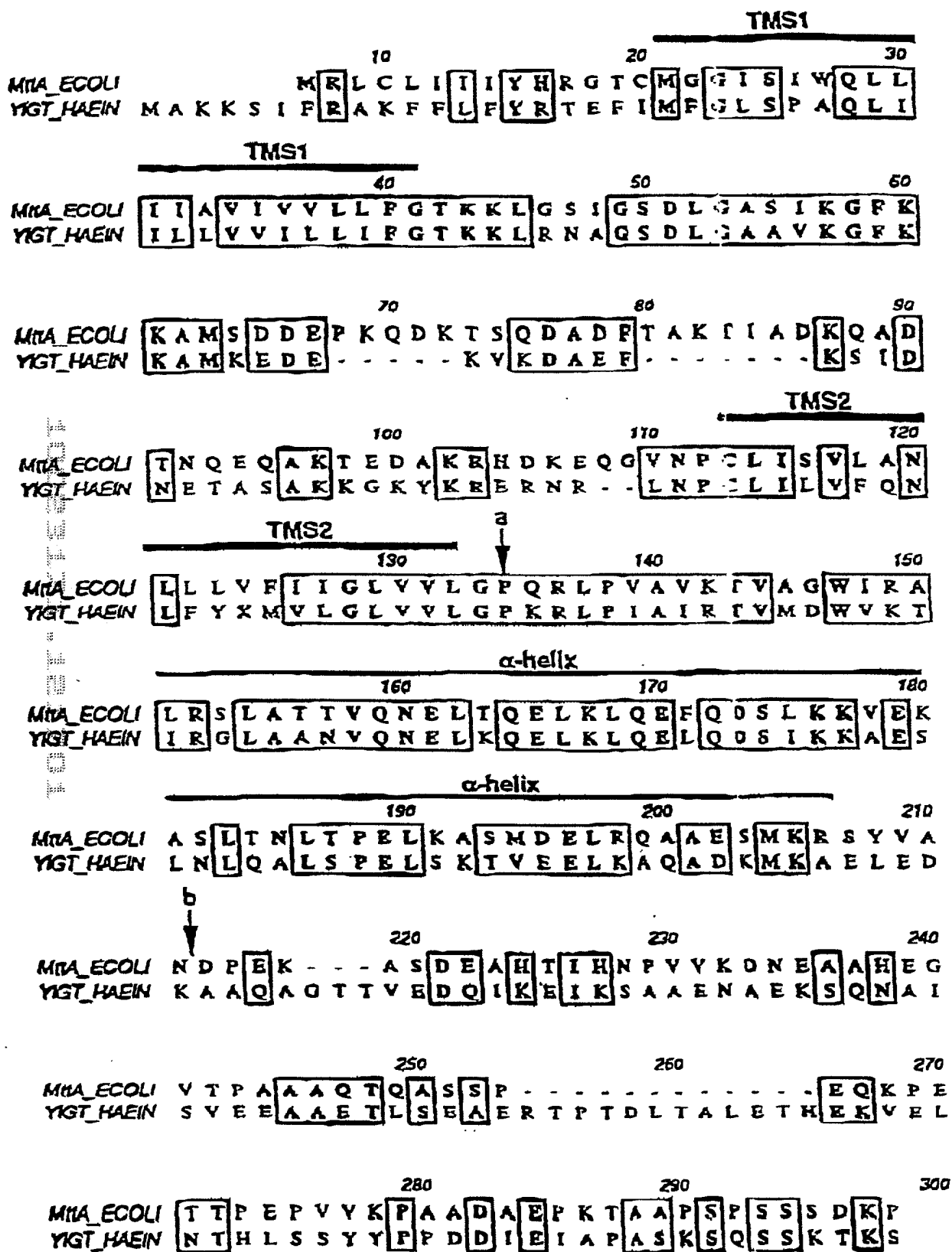


Figure 6



[illegible]

```

3000      3010      3020      3030      3040      3050
TCTGGCTGGGTGCCACCAGATACCAACCTTGAAGAGTTCGAATTTGCCATTCTGTACGGT

5060      5070      5080      5090      5100      5110
CTCTCAACCTATCTTTGAGAAACCCCTGGCCGAAATTTCTGTTTGGACATCTACTGTTAA

5120      5130      5140      5150      5160      5170
TCTGTTTAATACCGCGCGTCCCTTCAATATGCAAGTGCAGCCGCAACTCGTGTACTCCA

5180      5190      5200      5210      5220      5230
GAAACCCCTGCTCTACCTCCAAAGGOTAGGACGCCAGCTTTATCCGCAACTCGATTATG

5240      5250      5260      5270      5280      5290
GAAACCGGCGAAGCCTTTCTCTGAGTCTGTGATTAAAGATCAGGTGGGTATCTCTCCCT

5300      5310      5320      5330      5340      5350
GGTGAGACCATTTAAACAAAAGCCCGCTTCTGGGTCCAAAATAATCCCAACTGCTCTGA

5360      5370      5380      5390      5400      5410
ATTGGTTTACGACAGTTTCCGCCAGGGCAAGTATTACAGCACAGTGTGATAAGATTGC

5420      5430      5440      5450      5460      5470
CCCGGAGCTTCAGTCAAATCATGTACGTCAGGGACAATCCCGTATTTTCTCGGAATTGG

5480      5490      5500      5510      5520      5530
CGCTACGTTACTATTAACTGGCACATTCTTGTGGTCAGCCGACCTCAATGGGGGCTGAT

5540      5550      5560      5570      5580      5590
CCCCGGCTGGTTAATGGCGTGTGGTCTGATCGCTGCTTCTCGTTGGCGCAAAAGACG

5600      5610      5620      5630      5640      5650
CTGATTTTTCATCCCTCAAGCGCGCGCTGTAAAGTATAATGGCGCTTTGTTTAATCAT
                                     H R L C L I I>
                                     ____ORF RF[3] ____>

5660      5670      5680      5690      5700      5710
CATCTACCACAGAGGAACATGTATGGGTGGTATCAGTATTTGGCAGTTATTGATTATTGC
  I Y H R G T C M C G I S I W Q L L I I A>
  ____ORF RF[3] ____>

5720      5730      5740      5750      5760      5770
CGTCATCGTTGTACTGCTTTTGGCAACAAAAGCTCGCTCCATCGGTTCCGATCTTGG
  V I V V L L F G T K K L G S I G S D L G>
  ____ORF RF[3] ____>

5780      5790      5800      5810      5820      5830
TGGCTCCATCAAAGGCTTTAAAAAGCAATGAGCGATGATGAACCAAGCAGGATAAAC
  A S I K C F K K A M S D D E P K Q D K I>
  ____ORF RF[3] ____>

5840      5850      5860      5870      5880      5890
CAGTCAGGATGCTGATTTTACTGCGAAAACATACGCCGATACCAGCGCGATACGAATCA
  S Q D A D F T A K T I A D K Q A D T N Q>
  ____ORF RF[3] ____>

5900      5910      5920      5930      5940      5950

```

Figure 7(B)

GGAACAGGCTAAACAGAGACCGGAAGCGCCACGATAAAGAGCAGGTGAATCCGTGTTT
E Q A K T E D A K R H D K E Q V N F C L>
ORF RF(3) >

5960 5970 5980 5990 6000 6010
GATATCEGTTTTAGCGAACTTGGTATTGGTGTTCATCATCGGCTCGTCTCTCGGCGCC
I S V L A N L L L V F I I G L V V L C P>
ORF RF(3) >

6020 6030 6040 6050 6060 6070
GCAACCACTGCTGCGCGTAATAACCGTAGCGCGCTGGATTGCGCGCTTGGCTTCACT
Q R L F V A V K T V A G W I R A L R S L>
ORF RF(3) >

6080 6090 6100 6110 6120 6130
GGCGACAACGGTGCAGAACGAACTGACCCAGGACTTAAACTCCAGGAGTTTCAGGACAG
A T T V Q N E L T Q E L K L Q E F Q D S>
ORF RF(3) >

6140 6150 6160 6170 6180 6190
TCTGAAAAGGTTGAAAAGGCGACCCCTCACTAACCTGACGCGCGAACTGAAAAGCGTCGAT
L K K V E K A S L T N L T P E L K A S M>
ORF RF(3) >

6200 6210 6220 6230 6240 6250
GGATCAACTACGCCAGGTCGCGGAGTCGATGAAGCGTTCCTACGTTGCAAACGATCCTGA
D E L R Q A A E S M K R S Y V A N D P E>
ORF RF(3) >

6260 6270 6280 6290 6300 6310
AAAGGCGAGCGATGAGCGGCACACCATCCATAACCCGGTGGTGAAGATATGAAGCTCC
K A S D E A H T I H N P V V K D N E A A>
ORF RF(3) >

6320 6330 6340 6350 6360 6370
GCATCAGCGCGCTAACCGCTCGCGCTGCACAAAGCAGGCGAGTTCGCGGGAACAGAAAGCC
H E G V T P A A A Q T Q A S S P E Q K P>
ORF RF(3) >

6380 6390 6400 6410 6420 6430
AGAAACCAAGCGCAGCGCGCTGGTAAACCTGCTCGGAGCGTGAACCGAAACCGCTGC
E T T T E F V V K P A A D A E P K T A A>
ORF RF(3) >

6440 6450 6460 6470 6480 6490
ACCTTCCCTTCGTGGAGTGATAAACCGTAAACATGTCTGTAGAGATACTCAACCGCTT
H S V E D T Q P L>
ORF RF(2) >
P S P S S S D K P>
ORF RF(3) >

6500 6510 6520 6530 6540 6550
ATCAGCGCATCTGATTGAGCTCGCTAAGCGTCTGCTGAAGTCCATTATCGCGGTCATCGTG
I T H L I E L R K R L L H C I I A V I V>
ORF RF(2) >

6560 6570 6580 6590 6600 6610
ATATTCTGTGTCTGCTGTAATTCCGCAATGACAICTATCACCTGGTATCCGCGCCATTG

[illegible]

3

Figure 7(D)

7280 7290 7300 7310 7320 7330
 TGGAGTACAGGATGTTTCATATCGCCGTTAATTTGACCAGTTCCCAATTTGCCGAAAGACC
 K E Y R M F D I G V N L T S S Q F A K D>
 ORF RF(1) >

7340 7350 7360 7370 7380 7390
 CTCATGATGTTGTAGCGTCCGCTTTTCACCGCGGAGTTAATCGGCTACTCATCACCGGCA
 R D D V V A C A F D A G V N G L L I T G>
 ORF RF(1) >

7400 7410 7420 7430 7440 7450
 CTAACTCGCTCAAAAGCCAGCAGGCCCAAAAGCTGGCGGTCAGTATTCGTCCTGTGGT
 T N L R E S Q Q A Q K L A R Q Y S S C W>
 ORF RF(1) >

7460 7470 7480 7490 7500 7510
 CAAAGCGCGCGGTACATCCTCAGCAGCAGCCAGTCCCAAGCTGCCACTGAAGAAGCGA
 S T A G V H P R D S S Q W Q A A T E E A>
 ORF RF(1) >

7520 7530 7540 7550 7560 7570
 TTATTCAGCTGGCGCGGAGCCAGAACTGGTGGCGATGGTGAAATGTGGTCTCGACTTTA
 I I E L A A Q P E V V A I G E C G L D F>
 ORF RF(1) >

7580 7590 7600 7610 7620 7630
 ACCGCAACTTTTCGACGCCGGAAGAGCAGGAACGGGCTTTTGTGCCCCAGCTACGGATTG
 N R H F S T P E E Q E R A F V A Q L R I>
 ORF RF(1) >

7640 7650 7660 7670 7680 7690
 CCGCAGATTTAAACATGCCCGATTTATGCACTGTCCGGATGCCACAGAGCGGTTTATGA
 A A D L N H P V F M H C R D A H E R F H>
 ORF RF(1) >

7700 7710 7720 7730 7740 7750
 CATTCCTGGAGCCGTTGGCTGGATAACTGCGCTTCGCGCTTCTTCATTGCTTTACCGGCA
 T L L E P W L D K L P G A V L E C F T G>
 ORF RF(1) >

7760 7770 7780 7790 7800 7810
 CACGCGAAGAGATGCAGCCCTCCGTTGGCGCATGGAAATTTATATCGGCATTACCGGTTGG
 T R E E M Q A C V A B G I Y I G I T G W>
 ORF RF(1) >

7820 7830 7840 7850 7860 7870
 TTTCCGATGAACGACGCGGACTGGAGCTGCGGGAATTTTCCCGTTGATTCCCGCGGAAA
 V C D E R R G L E L R E L L P L I P A E>
 ORF RF(1) >

7880 7890 7900 7910 7920 7930
 AATTACTGATCGAACTGATGCGCGTATCTGCTCCCTCGGATCTCACGCCAAAGCCAT
 K L L I E T D A P Y L L P R D L T P K E>
 ORF RF(1) >

7940 7950 7960 7970 7980 7990
 CATCCCGCGCAACGAGCCAGCCCATCTGCCCATATTTGCACGATATTCGGCACTGGC

[illegible][illegible]

Figure 8(A)

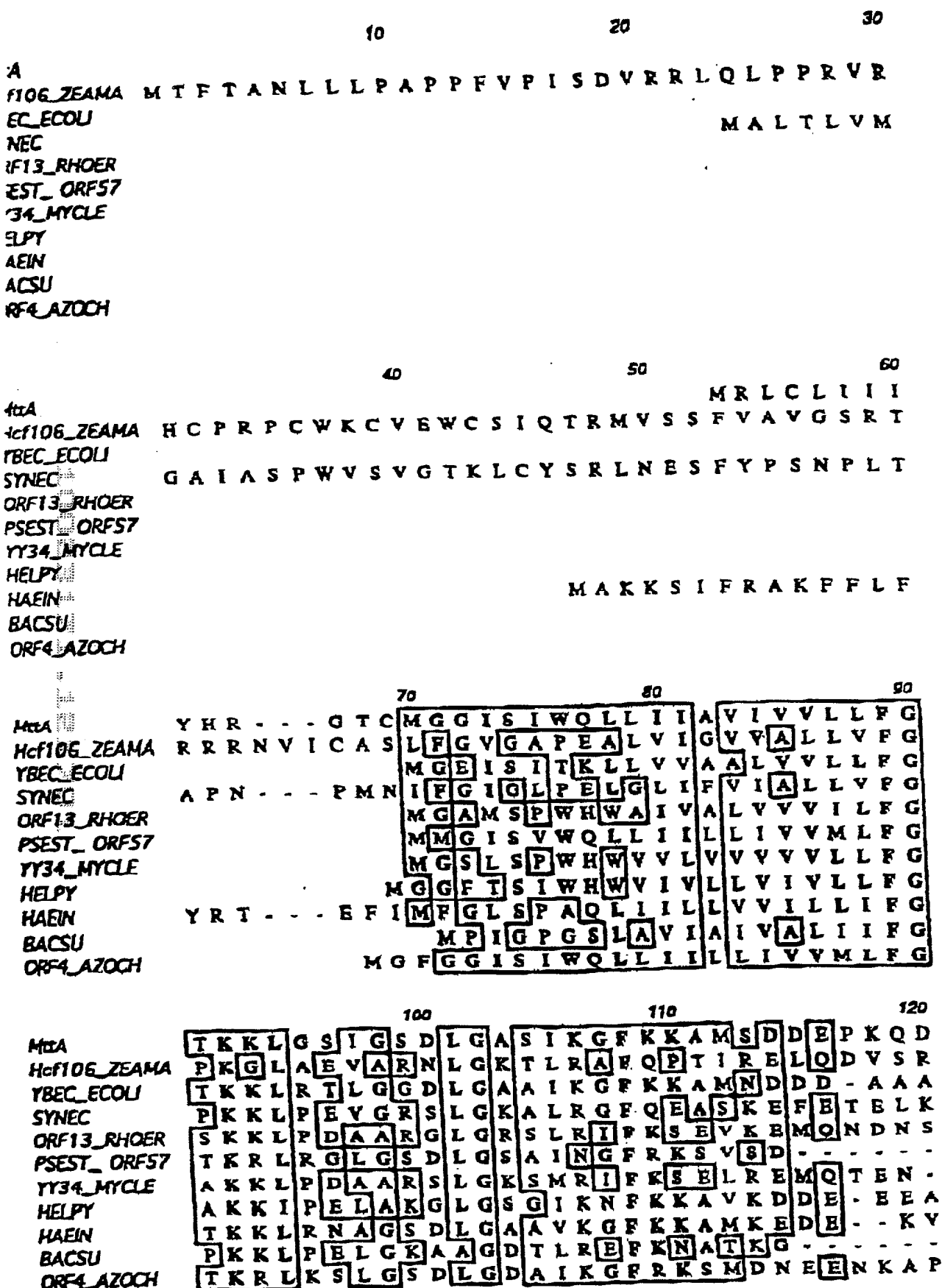


Figure 8(B)

	130	140	150
MtA	K T S Q D - A - - - D F T A K T I A D K Q A D T N Q E Q A K		
Hcf106_ZEAMA	E F R S T L E R E I G I D E V S Q S T K Y R P T T M N N N Q		
YBEC_ECOLI	K K G A D - V - - - D L Q A E K L S H K E		
SYNEC	R E A Q N L E - - - K S V Q I K A E L E E S K T P E S S S S		
ORF13_RHOER	T P A P T A Q - - - S A P P P Q S A P A E L P V A D T T T A		
PSEST_ORF57	- -		
YY34_MYCLE	- - - - - Q - - - - A Q A S A L E T P M Q N P T V V Q S Q R		
HELPY	K N E P - - K - - - T L D A Q A T Q T K V H E S S E I K S K		
HAEIN	K D A E F - K - - - S I D N E T A S A K K G K Y K R B R N R		
BACSU	- -		
ORF4_AZOCH	P V E E Q - K - - - G Q D H R G P G P Q G R G T G Q E R L S		

	160	170	180
MtA	T E D A K R H D K E Q G V N P C L I S V L A N L L L V F I I		
Hcf106_ZEAMA	Q -		
YBEC_ECOLI	- -		
SYNEC	- -		
ORF13_RHOER	P -		
PSEST_ORF57	- -		
YY34_MYCLE	- -		
HELPY	- -		
HAEIN	- -		
BACSU	- -		
ORF4_AZOCH	M F D I G -		F S E L L L V G L V

	190	200	210
MtA	G L V V L G P Q R L P V A V K T V A G W I R A L R S L A T T		
Hcf106_ZEAMA	- -		P A A D P N V K P E R A P
YBEC_ECOLI	- -		
SYNEC	- -		V T P P A P V
ORF13_RHOER	- -		S R S
PSEST_ORF57	- -		V V P P W S T
YY34_MYCLE	- -		
HELPY	- -		L N P C L I L
HAEIN	- -		
BACSU	- -		
ORF4_AZOCH	A L L V L G P E R L P V A A R M A G L W I G R L K R S F N T		

	220	230	240
MtA	V Q N E L T Q B L K L Q E F Q D S L K K V E K A S L T N L T		
Hcf106_ZEAMA	Y T S E E L M K V T E E Q I A A S A A A A W N P Q Q R A T S		
YBEC_ECOLI	- -		
SYNEC	- - - - - S E K A S		
ORF13_RHOER	Q P Q S Q H T E P K S A		
PSEST_ORF57	- -		
YY34_MYCLE	- - - - - E Q D H T E A R P A		
HELPY	- - - - - Q E S		

Figure 8(C)

AEIN VFQNLFY
 ACSU
 RF4_AZOCH LKT[**E**]VEREIGADEIRR...QLHNBRI BLE

260 260 270
 PELKASMDLRQAAESMKRSYVANDPEKAS
 HcF106_ZEAMA QQQEEAPTTFR - SEDAPTSGGSSGPAAPAR
 YBEC_ECOLI
 SYNEC
 ORF13_RHOER
 PSEST_ORF57
 YY34_MYCLE
 HELPY
 HAEIN
 BACSU
 ORF4_AZOCH REMKQSLQPPAPSA PDETAASVATPPQPAS

280 290 300
 DEAH TIHN PVVKDNEAAHEGVT PAAAQTQA
 HcF106_ZEAMA AESDSDPNQV NKSQKAEG ER
 YBEC_ECOLI
 SYNEC
 ORF13_RHOER
 PSEST_ORF57
 YY34_MYCLE
 HELPY
 HAEIN
 BACSU
 ORF4_AZOCH PAAHS DKTPSF

310 320 330
 SSPEQKPETTPEPVVKPAADAGPKTAAPSP
 HcF106_ZEAMA
 YBEC_ECOLI
 SYNEC
 ORF13_RHOER
 PSEST_ORF57
 YY34_MYCLE
 HELPY
 HAEIN
 BACSU
 ORF4_AZOCH

340 350 360
 SSSDKP
 HcF106_ZEAMA
 YBEC_ECOLI
 SYNEC
 ORF13_RHOER
 PSEST_ORF57
 YY34_MYCLE
 HELPY

Figure 9

MttB_ECOLI	I	T	H	L	I	E	L	R	K	R	L	L	N	C	I	I	A	V	I	V	I	-	F	L	C	L	V	Y	F	A	38
YC43_PORPU	T	E	H	L	E	E	L	R	Q	R	T	V	F	V	F	I	F	F	L	L	A	-	A	T	I	S	F	T	Q	I	58
YM16_MARPO	K	T	I	L	E	E	V	R	I	R	V	F	W	I	L	I	C	F	S	F	T	-	W	F	T	C	Y	W	F	S	34
ARATH	B	T	I	L	G	E	V	R	I	R	S	V	R	I	L	I	G	L	G	L	T	-	W	F	T	C	Y	W	F	S	43
Ymf16_RECAM	L	T	H	L	Y	E	I	R	L	R	I	I	Y	L	L	Y	S	I	P	L	T	-	C	F	C	S	Y	Q	Y	K	36
Y194_SYNY3	F	D	H	L	D	E	L	R	T	R	I	F	L	S	L	G	A	V	L	V	G	-	V	V	A	C	F	I	F	V	58
YY33_MYCTU	V	D	H	L	T	E	L	R	T	R	L	L	I	S	L	A	A	I	L	V	T	T	I	F	G	F	V	W	Y	S	57
HELPY	-	-	H	L	Q	E	L	R	K	R	L	M	V	S	V	G	T	I	L	V	A	-	F	L	G	C	F	H	F	W	34
TigU_HAEIN	I	T	H	L	V	E	L	R	N	R	L	L	R	C	V	I	C	V	V	L	V	-	F	V	A	L	V	Y	F	S	39
YcbT_BACSU	L	E	H	I	A	E	L	R	K	R	L	L	I	V	A	L	A	F	V	V	F	-	F	I	A	G	F	F	L	A	40
YH25_AZOCH	Y	A	H	L	T	E	L	R	S	B	L	L	R	S	V	A	A	V	L	L	I	-	F	A	A	L	F	Y	F	A	32
ARCFU	I	A	L	I	V	I	V	V	S	S	L	F	F	T	F	O	A	N	I	V	V	G	K	I	I	G	D	L	F	P	49

MttB_ECOLI	T	D	V	A	S	P	F	F	T	P	I	K	L	T	F	M	V	S	L	I	L	S	A	P	V	I	L	Y	Q	V	91
YC43_PORPU	L	A	P	G	E	Y	E	F	S	S	I	K	I	A	I	Y	C	G	I	V	A	T	T	P	F	G	V	Y	Q	V	106
YM16_MARPO	T	Q	L	T	E	A	L	S	T	Y	V	T	T	S	L	I	S	C	F	Y	F	L	F	P	E	L	S	Y	Q	I	87
ARATH	T	Q	L	T	E	A	F	S	T	F	V	A	T	S	S	I	A	C	S	Y	F	V	R	P	L	I	S	Y	Q	I	95
Ymf16_RECAM	T	D	L	I	E	A	F	I	T	Y	I	K	L	S	I	I	V	G	T	Y	L	S	Y	P	I	F	L	Y	Q	I	83
Y194_SYNY3	L	S	P	G	E	F	F	F	V	S	V	K	V	A	G	Y	S	G	I	L	V	M	S	P	F	I	L	Y	Q	I	106
YY33_MYCTU	T	A	P	F	D	Q	F	M	L	R	L	K	V	G	M	A	A	G	I	V	L	A	C	P	V	W	F	Y	Q	L	125
HELPY	L	S	P	I	E	G	V	M	V	A	V	K	I	S	F	S	A	A	I	V	I	S	M	P	I	I	F	W	Q	L	81
TigU_HAEIN	T	N	I	Q	T	P	F	F	T	P	I	K	L	T	A	I	V	A	I	F	I	S	V	P	Y	L	L	Y	Q	I	92
YcbT_BACSU	F	N	L	T	D	P	L	Y	V	F	M	Q	F	A	F	I	I	G	I	V	L	T	S	F	V	I	L	Y	Q	L	90
YH25_AZOCH	T	G	V	A	S	P	E	L	A	P	F	K	L	T	L	M	I	S	L	F	L	A	M	P	V	V	L	H	Q	V	85
ARCFU	L	T	P	L	E	G	L	L	L	Y	L	K	I	S	L	A	V	G	I	A	A	A	L	P	Y	I	F	H	L	V	139

MttB_ECOLI	W	A	F	I	A	P	-	-	-	A	L	Y	K	H	E	R	R	L	V	V	P	L	L	V	S	S	S	L	L	F	118
YC43_PORPU	I	L	Y	I	L	P	-	-	-	G	L	T	N	K	E	R	K	V	I	L	P	I	L	I	G	S	I	V	L	F	133
YM16_MARPO	W	C	F	L	M	P	-	-	-	S	C	Y	E	E	Q	R	K	K	Y	N	K	L	F	Y	L	S	G	F	C	F	114
ARATH	W	C	F	L	I	P	-	-	-	S	C	Y	G	E	Q	R	T	K	Y	N	R	F	F	Y	L	S	G	F	C	F	122
Ymf16_RECAM	W	S	F	L	I	P	-	-	-	G	F	P	L	Y	E	K	K	L	F	R	L	L	C	L	T	S	I	F	L	Y	110
Y194_SYNY3	I	Q	F	V	L	P	-	-	-	G	L	T	R	R	B	R	R	L	L	G	P	V	V	L	G	S	S	V	L	F	133
YY33_MYCTU	W	A	F	I	T	P	-	-	-	G	L	Y	Q	R	E	R	R	F	A	V	A	F	V	I	P	A	A	V	L	F	152
HELPY	W	L	F	I	A	P	-	-	-	G	L	Y	K	N	E	K	K	V	I	L	P	F	V	F	F	G	S	G	M	F	108
TigU_HAEIN	W	A	F	I	A	P	-	-	-	A	L	Y	Q	H	E	K	R	M	I	Y	P	L	L	P	S	S	T	I	L	F	119
YcbT_BACSU	W	A	F	V	S	P	-	-	-	G	L	Y	E	K	E	R	K	V	T	L	S	Y	I	P	V	S	I	L	L	F	117
YH25_AZOCH	W	G	F	I	A	P	-	-	-	G	L	Y	Q	H	E	K	R	I	A	M	P	L	M	A	S	S	V	L	L	F	112
ARCFU	L	T	A	L	R	E	R	G	V	I	T	P	S	F	R	K	T	S	A	P	K	Y	G	M	A	A	I	F	L	F	169

MttB_ECOLI	E	G	V	Q	V	S	T	D	I	A	S	Y	L	S	F	V	M	A	L	F	M	A	E	G	V	S	F	E	V	P	172
YC43_PORPU	D	I	V	E	P	L	W	S	F	E	Q	Y	F	D	F	I	L	L	L	F	S	T	G	L	A	F	E	I	P	187	
YM16_MARPO	L	I	I	K	L	Q	P	K	I	F	D	Y	I	M	L	T	V	R	I	L	F	I	S	S	I	C	S	Q	V	P	173
arab thal mito	L	M	I	K	L	Q	P	K	I	Y	D	Y	I	M	L	T	V	R	I	S	F	I	S	S	V	C	S	Q	V	P	181
Ymf16_RECAM	F	T	I	E	L	Q	A	K	I	H	E	Y	L	I	L	N	T	K	L	I	F	S	L	S	I	C	F	Q	L	P	170
Y194_SYNY3	D	V	V	E	Q	L	W	S	I	D	K	Y	F	E	F	V	L	L	L	M	F	S	T	G	L	A	F	Q	I	P	187
YY33_MYCTU	D	V	Q	V	T	A	L	S	G	D	R	Y	F	G	F	L	L	N	L	L	V	V	F	G	V	A	F	E	L	P	206
HELPY	D	V	F	A	A	N	I	S	A	S	S	Y	V	S	F	F	T	R	L	I	L	G	F	G	V	A	F	E	L	P	162
TigU_HAEIN	E	G	V	T	I	A	T	D	I	S	S	Y	L	D	F	A	L	A	L	F	L	A	F	G	V	C	F	E	V	P	173
YcbT_BACSU	L	N	V	N	Q	V	I	G	I	N	E	Y	F	H	F	L	L	Q	L	T	I	P	F	G	L	L	F	Q	M	P	171
YH25_AZOCH	E	G	V	A	M	M	T	D	I	G	Q	Y	L	D	F	V	L	T	L	F	P	A	F	G	V	A	F	E	V	P	160
ARCFU	Q	G	A	I	P	L	Y	S	L	S	E	F	V	N	F	V	A	L	M	L	V	L	E	G	I	V	F	E	L	P	222

Figure 10

TEEA	I	I	E	L	A	A	Q	-	-	P	E	V	V	A	I	G	E	C	G	L	D	F	N	R	N	F	104	
DVED	L	R	R	L	A	A	E	-	-	E	G	V	V	A	L	G	E	T	G	L	D	Y	Y	Y	T	P	101	
SLEQ	L	Q	Q	A	L	B	R	R	P	A	K	V	V	A	V	G	E	I	G	L	D	L	F	G	D	D	106	
LIGE	V	V	S	Q	I	E	S	N	I	D	L	I	V	A	V	G	E	T	G	M	D	F	H	H	T	R	107	
AQAT	L	K	K	L	V	S	T	H	R	S	F	I	S	C	I	G	E	Y	G	F	D	Y	H	Y	T	K	105	
ARAB	L	B	R	L	V	A	H	-	-	P	R	V	V	A	V	G	E	T	G	I	D	M	Y	W	P	G	102	
DES	L	F	E	K	F	V	G	H	-	-	Q	K	C	V	A	I	G	E	C	G	L	D	Y	Y	R	L	P	98
DAER	L	L	R	L	A	Q	D	-	-	P	K	V	I	A	I	G	E	I	G	L	D	Y	Y	Y	S	A	104	
DLAW	I	K	E	L	S	A	H	-	-	E	K	V	V	A	I	G	E	M	G	L	D	Y	H	W	D	K	101	
-EAL	A	N	K	G	K	A	S	-	-	G	K	V	V	A	F	G	E	F	G	L	D	Y	D	R	L	H	79	
HISK	M	E	Q	F	F	V	E	H	E	R	D	I	C	V	G	E	C	G	L	D	H	T	I	S	Q	211		
QERN	L	L	Q	A	L	R	H	-	-	P	K	A	V	A	F	G	E	M	G	L	D	Y	S	Y	K	C	602	

HCRDA	H	E	R	F	M	T	L	L	E	P	W	L	D	K	L	P	G	-	A	V	L	H	C	F	T	G	T	162		
HTRDA	R	A	D	T	L	A	I	L	R	E	E	K	V	T	D	C	G	-	G	V	L	H	C	F	T	B	D	160		
HSR	R	T	H	D	K	L	A	M	H	L	K	R	H	D	L	P	R	T	G	-	V	V	H	G	F	S	G	S	162	
HARDA	E	E	R	A	L	B	T	V	L	E	Y	R	V	P	B	V	-	-	I	F	H	C	Y	G	G	S	164			
HVR	D	V	H	E	R	I	Y	E	V	L	K	R	-	L	K	P	K	Q	P	V	V	F	H	C	F	S	E	D	161	
HNR	Q	A	D	R	D	V	L	D	V	L	R	A	E	G	A	P	D	T	-	V	I	L	H	C	F	S	S	D	163	
HIRE	A	S	F	D	S	L	N	L	L	K	N	-	-	Y	P	K	A	F	G	V	L	H	C	F	N	A	D	159		
HTRS	A	G	D	D	T	I	A	M	L	R	Q	H	R	A	E	K	C	G	-	G	V	I	H	C	F	T	E	T	161	
HNR	D	A	T	E	D	V	V	T	I	L	K	E	E	G	A	E	A	V	G	-	G	I	M	H	C	F	T	G	S	158
HSR	N	A	E	N	D	F	F	A	I	L	E	K	Y	L	P	E	L	P	K	K	G	V	V	H	S	F	T	G	S	138
HSRS	A	A	R	T	I	E	I	L	L	E	C	H	V	A	P	D	Q	-	V	V	L	H	A	F	D	G	T	282		
HCRE	A	D	E	D	L	L	B	I	M	K	K	F	V	P	P	D	Y	K	-	I	H	R	H	C	F	T	G	S	660	

ERR	G	L	E	L	R	E	L	L	P	L	I	P	A	E	K	L	L	I	E	T	D	A	P	Y	L	L	P	213	
RN	-	A	E	Q	L	R	D	A	A	R	Y	V	P	L	D	R	L	L	V	E	T	D	S	P	Y	L	A	P	209
PR	-	A	S	K	T	R	D	V	I	A	K	L	P	L	A	S	L	L	L	E	T	D	A	P	D	M	P	L	213
S	-	-	E	H	H	M	E	L	V	R	A	I	P	L	E	G	M	L	T	E	T	D	S	P	Y	L	S	-	212
KN	-	A	K	N	L	Q	A	A	L	S	V	I	P	T	E	L	L	L	S	H	T	D	S	P	Y	L	A	P	217
RT	-	A	R	E	L	R	E	A	V	P	L	M	P	V	E	Q	L	L	V	E	T	D	A	P	Y	L	T	P	214
KN	-	A	K	R	L	V	B	I	L	P	K	I	P	K	N	R	L	L	E	T	D	S	P	Y	L	T	P	208	
KN	-	A	E	A	I	R	E	V	I	R	Y	V	P	M	E	R	L	L	V	E	T	D	S	P	Y	L	A	P	212
KN	-	A	K	K	P	K	E	V	V	K	E	I	P	N	D	R	L	L	I	E	T	D	C	P	F	L	T	P	209
T	-	-	E	E	N	L	E	V	V	R	A	I	P	L	E	K	M	L	L	E	T	D	A	P	W	C	E	V	187
S	-	-	E	E	T	T	Q	L	I	E	S	I	P	L	S	Q	L	L	L	E	T	D	S	P	A	L	G	-	330
SS	-	A	W	E	A	R	E	A	L	R	Q	I	P	L	E	R	I	I	V	E	T	D	A	P	Y	F	L	P	713

Year	1990	1991	1992	1993	1994	1995	1996	1997	1998	1999	2000	2001	2002	2003	2004	2005	2006	2007	2008	2009	2010	2011	2012	2013	2014	2015	2016	2017	2018	2019	2020	2021	2022	2023	2024	2025	2026	2027	2028	2029	2030	2031	2032	2033	2034	2035	2036	2037	2038	2039	2040	2041	2042	2043	2044	2045	2046	2047	2048	2049	2050	2051	2052	2053	2054	2055	2056	2057	2058	2059	2060	2061	2062	2063	2064	2065	2066	2067	2068	2069	2070	2071	2072	2073	2074	2075	2076	2077	2078	2079	2080	2081	2082	2083	2084	2085	2086	2087	2088	2089	2090	2091	2092	2093	2094	2095	2096	2097	2098	2099
1990	1991	1992	1993	1994	1995	1996	1997	1998	1999	2000	2001	2002	2003	2004	2005	2006	2007	2008	2009	2010	2011	2012	2013	2014	2015	2016	2017	2018	2019	2020	2021	2022	2023	2024	2025	2026	2027	2028	2029	2030	2031	2032	2033	2034	2035	2036	2037	2038	2039	2040	2041	2042	2043	2044	2045	2046	2047	2048	2049	2050	2051	2052	2053	2054	2055	2056	2057	2058	2059	2060	2061	2062	2063	2064	2065	2066	2067	2068	2069	2070	2071	2072	2073	2074	2075	2076	2077	2078	2079	2080	2081	2082	2083	2084	2085	2086	2087	2088	2089	2090	2091	2092	2093	2094	2095	2096	2097	2098	2099	

10 20 30 40 50 60
TCTTGGCTGGGTGCCACCAGATACCAACGTTGAAGAGTTCGAATTTGCCATTGGTACGG

70 80 90 100 110 120
TCTGTGAACCTATCTTTGAGAAACCGCTCGCCGAAATTTGTTGGACATGTACTGTITAA

130 140 150 160 170 180
ATCTGTTTAATACGGCGCGCTCGCTTCAATATGGAAGTCCAGCCGCAACTCGTGTACTCC

190 200 210 220 230 240
AGAAAACCCCTGCTCTACGTCGAAGGGGTAGGACGCCAGCTTTATCCCAACTCGATTAT

250 260 270 280 290 300
GGAAAACGGCGAAGCCCTTTCCTCGAGTCGTGGATTAAAGATCAGGTCGGTATTCCTCGGC

310 320 330 340 350 360
TGGTGCACGCAATTTAAAGAAAAAGCGCCGTTCTGGGTGCAAAAATGCCAGAACTGCCTG

370 380 390 400 410 420
AATTGCTTTACGACAGTTTTCGCCAGGGCAAGTATTTCACAGCACAGCTTTCATAAGATTG

430 440 450 460 470 480
CCCCGGAGCTTCAGTCAAAATCATGTACGTCAGGCACAATCGCGTTATTTTCTCGGAATTG

490 500 510 520 530 540
CGGCTACGTTAGTATTAAGTGGCAGATTCTTCTGGTCAGCCCACTGCAATGGGGGCTGA

550 560 570 580 590 600
TGCCCGGCTGGTTAATGGCAGGTGCTCTGATCGCCCTCGTTTGTGGTTGGCGCAAAACAC

610 620 630 640 650 660
GCTGATTTTTTCATCGCTCAAGCGCGCGCGCTCTAACGTATATGCGGCTTTGTTTAACTCA
M K L C L I >

670 680 690 700 710 720
TCAICTACCACAGAGGAACATGTATGGGTGGTATCAGTATTTGGCAGTTATGATTATTG
I I Y H R G T C M C C I S I W Q L L I I >

730 740 750 760 770 780
CCGTCATCGTTGTACTGCTTTTTTGGCACCAAAAACCTCGGCTCCATCGGTTCCGATCTTG
A V I V V L L F G T K K L G S I G S D L >

790 800 810 820 830 840
GTGCGTCCATCAAAGGCTTTAAAAAAGCAATGAGCGATCATGACCAAAGCAGGATAAAA
G A S I K G F K K A M S D D E P K Q D K >

850 860 870 880 890 900
CCAGTCAGGATGCTGATTTTACTGCGAAAACTATCGCCGATAAGCAGCGCGGATACGGAATC
T S Q D A D F T A K T I A D K Q A D T N >

[illegible]

M S V E D T Q P L>

[illegible]

2110 2120 2130 2140 2150 2160
GCCCGATGCTTCTCGCAAAACGCTGTTGGCGATCCCGATGTACGTCTGTTTGAATCGG
P D V F S Q T L L A I P M Y C L F E I G>

2170 2180 2190 2200 2210 2220
TGTCTTCTTCTCAGCTTTTACGTTGGTAAAGGGCGAATCGGGACAGGAAACGACGC
V F F S R F Y V G K G R N R E E E N D A

[illegible]

2230 2240 2250 2260 2270 2280
TGAAGCAGAAAGCGAAAAAACTGAAGAATAAAATTC AACCGCCCGCTCAGGCCGGTTGTCAT
E A E S E K T E D>

2290 2300 2310 2320 2330 2340
ATGGACTACAGGATGTTTGATATCGGCGTTAATTIGACCAGTTCGCAATTTCCGAAGAC
M E Y R M F D I G V N L T S S Q F A K D>

2350 2360 2370 2380 2390 2400
CGTGATGATGTTGTAGCGTGC GCTTTTGACGCGGGAGTTAAIGGGTACTCATCACCGGC
R D D V V A C A F D A C V N G L L I T G>

2410 2420 2430 2440 2450 2460
ACTAACCTGCGTGAAAGCCACACGCGGCAAAAGCTGGCGCGTCAGTATTCGTCTCTGTGG
T N L R E S Q Q A Q K L A R Q Y S S C W>

2470 2480 2490 2500 2510 2520
TCAACGGCGGGCGTACATCCTCACCACAGCAGCCAGTGGCAAGCTGCGACTGAAGAAGCG
S I A C V H F H D S S Q W Q A A T E E A>

2530 2540 2550 2560 2570 2580
ATTATTGAGCTGCCCGCGCAGCCAGAAAGTGGTGGCGAATGGTGAATGTTGTTCTCGACTTT
I I E L A A Q P E V V A I G E C G L D F>

2590 2600 2610 2620 2630 2640
AACC CGAACTTTTCGACGCGCGGAAGAGCAGGAACGCGCTTTTGTGGCCAGCTACGCATT
N R N F S T P E E Q E R A F V A Q L R I>

2650 2660 2670 2680 2690 2700
GCCGCAGATTTAAACATGCCGGTATTTATGCACTGTGCGGATGCCACGACGGGTTTATG
A A D L N M P V F M H C R D A H E R F M>

2710 2720 2730 2740 2750 2760
ACATTGCTGGAGCCCTCCCTGGATAAACTGCCCTGGTGGCGTTCTTCATTGCTTTACCGGC
T L L E P W L D K L P G A V L H C F T C>

2770 2780 2790 2800 2810 2820
ACACCCGAAGAGATGCACCCGTGCGTGGCCCGAATTTATATCGGCATTACCGGTTGG
T R E E M Q A C V A H G I Y I G I T G W>

2830 2840 2850 2860 2870 2880
GTTTGGGATGAACGACGCCGACTGGACCTGCCCGAATTTTGGCGTTGATTCCGGCGGAA
V C D E R R G L E L R E L L P L I F A D>

Country	Year	Population (millions)	Urban population (millions)	Urban population (%)	Population density (per sq km)	Urban population density (per sq km)
Algeria	1980	11.0	5.0	45.5	10.0	10.0
Algeria	1985	11.5	5.5	47.8	10.5	10.5
Algeria	1990	12.0	6.0	50.0	11.0	11.0
Algeria	1995	12.5	6.5	52.0	11.5	11.5
Algeria	2000	13.0	7.0	53.8	12.0	12.0
Algeria	2005	13.5	7.5	55.6	12.5	12.5
Algeria	2010	14.0	8.0	57.1	13.0	13.0
Algeria	2015	14.5	8.5	58.6	13.5	13.5
Algeria	2020	15.0	9.0	60.0	14.0	14.0
Algeria	2025	15.5	9.5	61.3	14.5	14.5
Algeria	2030	16.0	10.0	62.5	15.0	15.0
Algeria	2035	16.5	10.5	63.6	15.5	15.5
Algeria	2040	17.0	11.0	64.7	16.0	16.0
Algeria	2045	17.5	11.5	65.7	16.5	16.5
Algeria	2050	18.0	12.0	66.7	17.0	17.0
Algeria	2055	18.5	12.5	67.6	17.5	17.5
Algeria	2060	19.0	13.0	68.4	18.0	18.0
Algeria	2065	19.5	13.5	69.2	18.5	18.5
Algeria	2070	20.0	14.0	70.0	19.0	19.0
Algeria	2075	20.5	14.5	70.7	19.5	19.5
Algeria	2080	21.0	15.0	71.4	20.0	20.0
Algeria	2085	21.5	15.5	72.1	20.5	20.5
Algeria	2090	22.0	16.0	72.7	21.0	21.0
Algeria	2095	22.5	16.5	73.3	21.5	21.5
Algeria	2100	23.0	17.0	73.9	22.0	22.0
Algeria	2105	23.5	17.5	74.5	22.5	22.5
Algeria	2110	24.0	18.0	75.0	23.0	23.0
Algeria	2115	24.5	18.5	75.5	23.5	23.5
Algeria	2120	25.0	19.0	76.0	24.0	24.0
Algeria	2125	25.5	19.5	76.5	24.5	24.5
Algeria	2130	26.0	20.0	76.9	25.0	25.0
Algeria	2135	26.5	20.5	77.3	25.5	25.5
Algeria	2140	27.0	21.0	77.8	26.0	26.0
Algeria	2145	27.5	21.5	78.2	26.5	26.5
Algeria	2150	28.0	22.0	78.6	27.0	27.0
Algeria	2155	28.5	22.5	79.0	27.5	27.5
Algeria	2160	29.0	23.0	79.3	28.0	28.0
Algeria	2165	29.5	23.5	79.7	28.5	28.5
Algeria	2170	30.0	24.0	80.0	29.0	29.0
Algeria	2175	30.5	24.5	80.3	29.5	29.5
Algeria	2180	31.0	25.0	80.6	30.0	30.0
Algeria	2185	31.5	25.5	81.0	30.5	30.5
Algeria	2190	32.0	26.0	81.3	31.0	31.0
Algeria	2195	32.5	26.5	81.6	31.5	31.5
Algeria	2200	33.0	27.0	81.8	32.0	32.0
Algeria	2205	33.5	27.5	82.1	32.5	32.5
Algeria	2210	34.0	28.0	82.4	33.0	33.0
Algeria	2215	34.5	28.5	82.6	33.5	33.5
Algeria	2220	35.0	29.0	82.9	34.0	34.0
Algeria	2225	35.5	29.5	83.1	34.5	34.5
Algeria	2230	36.0	30.0	83.3	35.0	35.0
Algeria	2235	36.5	30.5	83.6	35.5	35.5
Algeria	2240	37.0	31.0	83.8	36.0	

2890	2900	2910	2920	2930	2940
AAATTATCATCGAAACTGATCGCCCGTATCTGCTCCCTCGCGATCTCAAGCCAAAGCCA					
K L L I E T D A P Y L L P R D L T P K P >					
<hr/>					
2950	2960	2970	2980	2990	3000
TCATCCCCCGCGCAACCAACCCAGCCCATCTGCCCCATATTTTGCACGGTATTGCCCACTGG					
S S R R N E P A H L P H I L Q R I A M W >					
<hr/>					
3010	3020	3030	3040	3050	3060
CGTCGAGAAAGATGCGCGCATGGCTGGCTGCCACCAACGGATGCTAATGCCAAAACACTGTITT					
R G E D A A W L A A T T I D A N A K T L F >					
<hr/>					
3070	3080	3090	3100	3110	3120
GGGATTGCGTIIITAGAGITTTGCGGAACCTCGGTATTCTTCACACTGTGCTTAATCTCTTTA					
G I A F >					